Raw Sequence Listing Error Summary

| ERROR DETECTED | SUGGESTED CORRECTION | SERIAL NUMBER: _09/287,995_ |

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 _____ Wrapped Nucleics
Wrapped Aminos
The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 _____ Invalid Line Length
The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 _____ Misaligned Amino
Numbering
The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4 _____ Non-ASCII
The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5 _____ Variable Length
Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 _____ PatentIn 2.0
"bug"
A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7 _____ Skipped Sequences
(OLD RULES)
Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8 _____ Skipped Sequences
(NEW RULES)
Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 _____ Use of n's or Xaa's
(NEW RULES)
Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 _____ Invalid <213>
Response
Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 _____ Use of <220>
Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32)   (Sec. 1.823 of Sequence Rules)

12 _____ PatentIn 2.0
"bug"
Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 _____ Misuse of n
n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH – Biotechnology Systems Branch – 08/21/2001

PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/787,995

DATE: 01/22/2002
TIME: I6:33:33

Input Set : A:\ST98032.ST25.txt
Output Set: N:\CRF3\01182002\I787995.raw

Does Not Comply
Corrected Diskette Need:

```
 3 <110> APPLICANT: Aventis Pharma S.A.
 5 <120> TITLE OF INVENTION: Use Of Specific Hybrid Promoters For Controlling Tissue
Expression
 7 <130> FILE REFERENCE: ST98032
 9 <140> CURRENT APPLICATION NUMBER: US/09/787,995
 9 <141> CURRENT FILING DATE: 2001-03-23
 9 <150> PRIOR APPLICATION NUMBER: US 60/123298
10 <151> PRIOR FILING DATE: 1999-04-04
12 <150> PRIOR APPLICATION NUMBER: FR/98/12000
13 <151> PRIOR FILING DATE: 1998-09-25
15 <160> NUMBER OF SEQ ID NOS: 10
17 <170> SOFTWARE: PatentIn version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 23
21 <212> TYPE: DNA
22 <213> ORGANISM: Artificial
24 <220> FEATURE:
25 <223> OTHER INFORMATION: Artificial Sequence
27 <220> FEATURE:
28 <221> NAME/KEY: nucleotide
29 <222> LOCATION: (1)..(23)
31 <400> SEQUENCE: 1
32 gatggtccct acttatgctg cta                                    23
35 <210> SEQ ID NO: 2
36 <211> LENGTH: 23
37 <212> TYPE: DNA
38 <213> ORGANISM: Artificial
40 <220> FEATURE:
41 <223> OTHER INFORMATION: articifial sequence
43 <220> FEATURE:
44 <221> NAME/KEY: nucleotide
45 <222> LOCATION: (1)..(23)
47 <400> SEQUENCE: 2
48 cttccatcat accaaactac ata                                    23
51 <210> SEQ ID NO: 3
52 <211> LENGTH: 32
53 <212> TYPE: DNA
54 <213> ORGANISM: Artificial
56 <220> FEATURE:
57 <223> OTHER INFORMATION: artificial sequence
59 <220> FEATURE:
60 <221> NAME/KEY: nucleotide
61 <222> LOCATION: (1)..(32)
63 <400> SEQUENCE: 3
```

Handwritten annotations:
- give source of genetic material — see item //
- on Error Summary Sheet
- misspelled (artificial) — see item //
- (artificial sequence) — item //

```
                    RAW SEQUENCE LISTING                  DATE: 01/22/2002
                    PATENT APPLICATION:  US/09/787,995    TIME: 16:33:33

                    Input Set : A:\ST98032.ST25.txt
                    Output Set: N:\CRF3\01182002\I787995.raw

      64 ctgctaaatt gctcgaqgac aaattagaca aa                               32
      67 <210> SEQ ID NO: 4
      68 <211> LENGTH: 33
      69 <212> TYPE: DNA
C--> 70 <213> ORGANISM: Artificial
      72 <220> FEATURE:
      73 <223> OTHER INFORMATION: artificial sequence
      75 <220> FEATURE:
W--> 76 <221> NAME/KEY: oligonucleotide
      77 <222> LOCATION: (1)..(33)
      79 <400> SEQUENCE: 4
      80 ccctgacaaa gcttggctgg gctgctccac tgg                             33
      83 <210> SEQ ID NO: 5
      84 <211> LENGTH: 30
      85 <212> TYPE: DNA
C--> 86 <213> ORGANISM: Artificial
      88 <220> FEATURE:
      89 <223> OTHER INFORMATION: artificial sequence
      91 <220> FEATURE:
W--> 92 <221> NAME/KEY: oligonucleotide
      93 <222> LOCATION: (1)..(30)
      95 <400> SEQUENCE: 5
      96 atcgacgcgt gcccgttaca taacttacgg                                 30
      99 <210> SEQ ID NO: 6
     100 <211> LENGTH: 38
     101 <212> TYPE: DNA
C--> 102 <213> ORGANISM: Artificial
     104 <220> FEATURE:
     105 <223> OTHER INFORMATION: artificial sequence
     107 <220> FEATURE:
W--> 108 <221> NAME/KEY: oligonucleotide
     109 <222> LOCATION: (1)..(38)
     111 <400> SEQUENCE: 6
     112 atcgacgcgt ccgctcgagc gtcaatgggg cggagttg                        38
     115 <210> SEQ ID NO: 7
     116 <211> LENGTH: 36
     117 <212> TYPE: DNA
C--> 118 <213> ORGANISM: Artificial
     120 <220> FEATURE:
     121 <223> OTHER INFORMATION: artificial sequence
     123 <220> FEATURE:
W--> 124 <221> NAME/KEY: oligonucleotide
     125 <222> LOCATION: (1)..(36)
     127 <400> SEQUENCE: 7
     128 ccaggctgca ctcgagacta gttcccacca actcga                          36
     131 <210> SEQ ID NO: 8
     132 <211> LENGTH: 36
     133 <212> TYPE: DNA
C--> 134 <213> ORGANISM: Artificial
```

```
                    RAW SEQUENCE LISTING              DATE: 01/22/2002
                    PATENT APPLICATION:  US/09/787,995   TIME: 16:33:33

                    Input Set : A:\ST98032.ST25.txt
                    Output Set: N:\CRF3\01182002\I787995.raw

    136 <220> FEATURE:
    137 <223> OTHER INFORMATION: artificial sequence
    139 <220> FEATURE:
W--> 140 <221> NAME/KEY: oligonucleotide
    141 <222> LOCATION: (1)..(36)
    143 <400> SEQUENCE: 8
    144 tcgtttgaag cttggaagga gagtagcttc ggtgtc                      36
    147 <210> SEQ ID NO: 9
    148 <211> LENGTH: 30
    149 <212> TYPE: DNA
C--> 150 <213> ORGANISM: Artificial
    152 <220> FEATURE:
    153 <223> OTHER INFORMATION: artificial sequence
    155 <220> FEATURE:
W--> 156 <221> NAME/KEY: oligonucleotide
    157 <222> LOCATION: (1)..(30)
    159 <400> SEQUENCE: 9
    160 cccgttacat aacttacggt aaatggcccg                             30
    163 <210> SEQ ID NO: 10
    164 <211> LENGTH: 30
    165 <212> TYPE: DNA
C--> 166 <213> ORGANISM: Artificial
    168 <220> FEATURE:
    169 <223> OTHER INFORMATION: artificial sequence
    171 <220> FEATURE:
W--> 172 <221> NAME/KEY: oligonucleotide
    173 <222> LOCATION: (1)..(30)
    175 <400> SEQUENCE: 10
    176 gggacgcgct tctacaaggc gctggccgaa                             30
```

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/787,995

DATE: 01/22/2002
TIME: 16:33:34

Input Set : A:\ST98032.ST25.txt
Output Set: N:\CRF3\01182002\I787995.raw

```
L:9   M:270 C: Current Application Number differs, Replaced Current Application No
L:9   M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:22  M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:28  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:38  M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:44  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:54  M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:60  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:70  M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:4
L:76  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:86  M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:92  M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:102 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:108 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:118 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:124 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:7
L:134 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:140 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:8
L:150 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:156 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:9
L:166 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:172 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10
```

ψK

BIOTECHNOLOGY
SYSTEMS
BRANCH

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information
Center (STIC) detected errors when processing the following computer readable
form:

Application Serial Number: _O9/787,995_
Source: _____ _PCT/09_
Date Processed by STIC: _____ _1/22/2002_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE
   APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
   NOTICE TO COMPLY
FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE: SEE BELOW FOR ADDRESS:
        http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is
a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom, including:
1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission
   User Manual - ePAVE)
2. U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
   U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
   Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
            Or
   U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
   2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office,
   Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202